

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:01:55 ; Search time 109 Seconds  
 (without alignments)  
 20559.273 Million cell

updates/sec

Title: US-09-300-482-225  
 Perfect score: 316  
 Sequence: 1  
 gataaggtgcgacactgag.....gaatggaggaggagtacttcga 316

Scoring table: IDENTITY\_NUC  
 Gapop 10.0 , Gapext 1.0

Searched: 9875436 seqs, 3558593875 residues

Total number of hits satisfying chosen parameters: 19750872

Minimum DB seq length: 0  
 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
 Maximum Match 100%  
 Listing first 45 summaries

Database : Issued\_Patents\_NA:\*  
 1: /ABSS/Data/CRF/ptodata/1/ina/1\_COMB.seq:\*  
 2: /ABSS/Data/CRF/ptodata/1/ina/5\_COMB.seq:\*  
 3: /ABSS/Data/CRF/ptodata/1/ina/6A\_COMB.seq:\*  
 4: /ABSS/Data/CRF/ptodata/1/ina/6B\_COMB.seq:\*  
 5: /ABSS/Data/CRF/ptodata/1/ina/7A\_COMB.seq:\*  
 6: /ABSS/Data/CRF/ptodata/1/ina/7B\_COMB.seq:\*  
 7: /ABSS/Data/CRF/ptodata/1/ina/7C\_COMB.seq:\*  
 8: /ABSS/Data/CRF/ptodata/1/ina/7D\_COMB.seq:\*  
 9: /ABSS/Data/CRF/ptodata/1/ina/HA\_COMB.seq:\*  
 10: /ABSS/Data/CRF/ptodata/1/ina/HB\_COMB.seq:\*  
 11: /ABSS/Data/CRF/ptodata/1/ina/PCTUS\_COMB.seq:\*  
 12: /ABSS/Data/CRF/ptodata/1/ina/PP\_COMB.seq:\*  
 13: /ABSS/Data/CRF/ptodata/1/ina/RE\_COMB.seq:\*  
 14: /ABSS/Data/CRF/ptodata/1/ina/backfiles1.seq:\*